

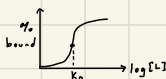
Enzyme Equations to know

$$K_D = \frac{[P][L]}{[P+L]} = \frac{1}{K_A}$$

↓ K_D = stronger protein-ligand binding

(K_i is just K_D for inhibitors specifically)

$K_D = [L]$ when 50% bound to Protein

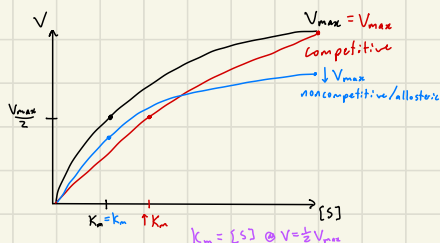


$$V_{max} = k_{cat} [E]$$

$$\text{catalytic efficiency} = \frac{k_{cat}}{K_m}$$

$$V_0 = \frac{V_{max} \times [S]}{K_m + [S]}$$

Enzyme Plots



so in summary:

competitive

= V_{max}

↑ K_m

noncompetitive/allosteric

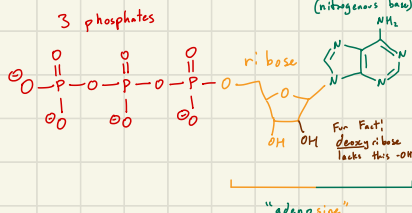
↓ V_{max}

= K_m

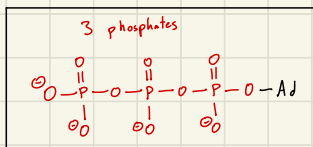
Good Inhibitors tend to 1) mimic substrate intermediates

2) mimic substrate peptide sequence

ATP



but for Chem 27 only need to know this abbreviation:



Names

ATP = ad - 3 phosphates

ADP = ad - 2 phosphates

AMP = ad - 1 phosphate

P_i = free PO_4^{3-}

PP_i = free $P_2O_7^{4-}$

Phosphoric Acid

relevant



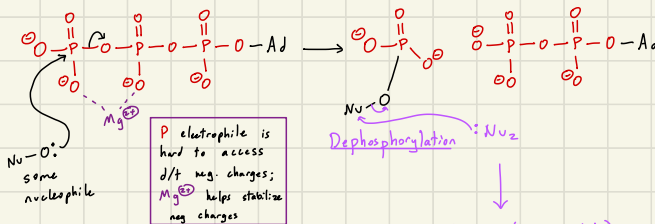
Fun Fact! H_3PO_4 $pK_{a1} = 2.1$, $pK_{a2} = 7.2$, $pK_{a3} = 12$

Phosphate Bonds

Thermodynamically favorable $\Delta G \ll 0$

Kinetically unfavorable d/t neg charges

(needs Mg^{2+} to lower E_a)



Dephosphorylation: Nu_2

P_i (inorganic free phosphate)

+ $Nu-Nu_2$